

Applicants: Prayaga et al.
U.S.S.N. 09/973,424



++ +e v g++t+y lt L+P ++Y ++ + + + S

SEQ ID NO: 68 468 gsITETLVQGDKTEYLLTALEPKSTYIICMV---TM--ETS 503

Please replace the pending sequence listing with the enclosed sequence listing, pages 1-59.

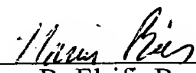
REMARKS

In response to the March 27, 2002 Notice to Comply with Requirements for Patent Applications Containing Nucleotide Sequence and/or Amino Acid Sequence Disclosures, Applicants enclose herein a substitute sequence listing. The specification has been amended to correct SEQ ID NOs in compliance with 37 C.F.R. §§ 1.821-1.825 and to include the substitute sequence listing. No new matter has been added.

Although Applicants believe no fees are due at this time, the Commissioner is hereby authorized to charge any fees that may be due, or credit any overpayment of same, to Deposit Account No. 50-0311, Attorney Reference No. 15966-585CIP2 (Cura-85CIP2). Should any questions or issues arise concerning this application, the Examiner is encouraged to contact the undersigned at the telephone number provided below.

Respectfully submitted,

April 25, 2002


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VERSION WITH MARKINGS TO SHOW CHANGES MADE

The paragraph beginning on page 37, line 21 has been amended as follows:

-- LRR: domain 9 of 9, from 285 to 308: score 28.1, E = 0.00021

->nLeeLdLsnN.LtSlppglfsnLp<- (SEQ ID NO: 73[85])

Le+LdLsnN+Lt+lp glf++L

SEQ ID NO: 68 285 ELERLDLSNNnLTTLPRGLFDDLg 308

LRRCT: domain 1 of 1, from 318 to 369: score 48.0, E = 2.1e-10

->NPfnCDCeLrwLlrWlretnprlledgedlrCasPeslrGqpIllellpsdfsCp<- (SEQ ID NO: 71[86])

NP+ C C+L+wL++W+++ ++ +++ + l+C++Pe++rG+ +++++ ++ *

SEQ ID NO: 68 318 NPWFCGCNLMWLRDWVKA-RAAVVNVR-GLMCQGPEKVRGMAIKDITSEVESVL 362

fn3: domain 1 of 1, from 425 to 503: score 13.8, E = 0.12

*->PsaPtntlvtvtdvtstsltlSwsptgngpitgYevtyRqpkngge..

+ ++v+++t++s++++W++ p ++++++ ++ +

SEQ ID NO: 68 425 GAKTLAIHVKALTADSIRITWKAT---LPASSFRLSW-LRLGHSPav 467

..wneltvpgtttsytltgLkPgteYevrVqAvnggGGpeS<-* (SEQ ID NO: 72[87])

++ +e v g++t+y lt L+P ++Y ++ + + S

SEQ ID NO: 68 468 gsITETLVQGDKTEYLLTALEPKSTYIICMV---TM--ETS 503 --